



**ARTICLE**

# DT-Net: Joint Dual-Input Transformer and CNN for Retinal Vessel Segmentation

Wenran Jia<sup>1</sup>, Simin Ma<sup>1</sup>, Peng Geng<sup>1</sup> and Yan Sun<sup>2,\*</sup>

<sup>1</sup>School of Information Science and Technology, Shijiazhuang Tiedao University, Shijiazhuang, China

<sup>2</sup>School of Mathematics and Information Science, Zhangjiakou University, Zhangjiakou, China

\*Corresponding Author: Yan Sun. Email: sunyan7322@163.com

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## ABSTRACT

Retinal vessel segmentation in fundus images plays an essential role in the screening, diagnosis, and treatment of many diseases. The acquired fundus images generally have the following problems: uneven illumination, high noise, and complex structure. It makes vessel segmentation very challenging. Previous methods of retinal vascular segmentation mainly use convolutional neural networks on U Network (U-Net) models, and they have many limitations and shortcomings, such as the loss of microvascular details at the end of the vessels. We address the limitations of convolution by introducing the transformer into retinal vessel segmentation. Therefore, we propose a hybrid method for retinal vessel segmentation based on modulated deformable convolution and the transformer, named DT-Net. Firstly, multi-scale image features are extracted by deformable convolution and multi-head self-attention (MHSA). Secondly, image information is recovered, and vessel morphology is refined by the proposed transformer decoder block. Finally, the local prediction results are obtained by the side output layer. The accuracy of the vessel segmentation is improved by the hybrid loss function. Experimental results show that our method obtains good segmentation performance on Specificity (SP), Sensitivity (SE), Accuracy (ACC), Curve (AUC), and F1-score on three publicly available fundus datasets such as DRIVE, STARE, and CHASE\_DB1.

## KEYWORDS

Retinal vessel segmentation; deformable convolution; multi-scale; transformer; hybrid loss function

## 1 Introduction

The vessels in fundus images are currently the only microvascular system that can be directly visualized non-invasively and painlessly. The pathological characteristics of related diseases can be observed by the morphology and changing information of retinal vessels. For example, diabetic patients are prone to retinopathy, macular degeneration, and blindness [1–3]. The retinal vessels of hypertensive patients have higher curvature and narrowing, which can easily lead to retinal hemorrhage [4]. Therefore, visualizing the distribution and details of retinal vessels can help doctors diagnose diseases more efficiently [5]. However, retinal vessels have the following problems: complex and diverse structures, tiny vessels, low contrast, and easy confusion with the background. A significant amount



of time and effort is required to segment the vessels manually. Therefore, an automatic retinal vessel segmentation method is essential to assist doctors in diagnosing diseases quickly.

Artificial intelligence technology shortens the distance between human life and computer, and the method based on deep learning is applied to various tasks. For example, Sultan et al. used deep learning to deal with the segmentation task of high-resolution aerial images [6,7]. Liu et al. [8] and Qin et al. [9] applied deep learning to image fusion. Jin et al. [10] applied deep learning to classification tasks to provide accurate ERM automatic grading for clinical practice. Deep learning is also widely used in the task of medical image segmentation [11,12] and other fields [13,14]. Among them, convolutional neural networks (CNN) have made great progress in location-sensitive tasks [15–17]. In recent years, U-Net [18] based on the full convolution network (FCN) [19] has been widely used in medical image segmentation [20–24]. However, U-Net is difficult to deal with irregular and tiny vessels. M-Net [25] is an improved U-Net framework that uses the image pyramid mechanism to realize multi-level receptive fields and can learn image features at different scales. However, the feature filtering is not realized in the hopping connection in the M-Net model. ResU-Net [26] is derived from the U-Net architecture. It uses residual blocks to replace convolutional layers and increases the depth of the model to get more vessel features. But the contrast-limited adaptive histogram equalization (CLAHE) operation increases the noise of the image. UNet++ [27] redesigns the skip connection part of U-Net, which is the aggregation of features of different semantic scales in the decoder. But it consumes too much memory and takes a lot of time on small datasets. IterNet [28] is an encoder-decoder model like U-Net, which adopts U-Net as the basic module to improve the connectivity of vessel segmentation results by expanding the depth of the model through multiple iterations.

Based on U-Net, Deformable U-Net (DUNet) [29] adds deformable convolution [30] to adaptively adjust the receptive field according to the size and shape of vessels and improve segmentation accuracy and noise immunity. MAU-Net [31] uses modulated deformable convolution [32] as an encoding and decoding unit and uses position and channel attention block to realize vascular segmentation. Recently, the transformer has been successfully applied to the field of computer vision. Inspired by this, TransUNet uses a hybrid of CNN and transformer as an encoder and uses a skip connection and decoder for medical image segmentation [33]. The encoder, bottleneck, and decoder of Swin-Unet [34] use the Swin-transformer block [35] to realize medical image segmentation. FAT-Net [36] implements a dual encoder, including both CNN and transformer branches, to achieve skin lesion segmentation. Although it can get better performance, these models based on the transformer are both complicated and time-consuming, which will influence the practicability to some extent.

These segmentation methods have the following problems: (1) The method used can only extract the local information in the image and can not deal with the global features. (2) The accuracy of segmentation is low. (3) The structural information in the vascular image can not be obtained well. Given the above problems, we use deformable convolution to extract complex and variable structural information, which has better learning ability than ordinary convolution. In addition, we use the transformer to capture long-term dependencies through a self-attention mechanism and help CNN overcome its inherent spatially induced biases [37]. Therefore, a segmentation network based on a combination of deformable convolution [32] and transformer is proposed to solve the challenging task of retinal vessel segmentation. The proposed network uses convolution to extract local features and the transformer to construct long-term dependencies. And pre-training on large-scale datasets is not required and achieves better results on small-scale datasets. Our main contributions are summarized as follows:

(1) We propose an end-to-end deep learning network named DT-Net, which is very effective for retinal vessel segmentation. The network takes into account multi-scale input, structural information, and long-term dependency, and provides more powerful technical support for clinical diagnosis and processing.

(2) Combine deformable convolution with transformer. Deformable convolution can extract structural information in retinal vessels. Transformer makes up for the defect that CNN can only obtain local information and enhances the extraction ability of feature information to achieve a better segmentation effect.

(3) A dual-input MHSA algorithm was proposed to extract multi-scale image information of fundus vascular images with different resolutions. The output of multi-scale image information is fused by skip connection to compensate for the information loss in feature extraction. A mixed loss function was used to improve the accuracy of retinal vessel segmentation.

(4) We conducted experiments on DRIVE, STARE, and CHASE\_DB1 with accuracy rates of 96.31%, 97.03%, and 97.37%, respectively. The experimental results showed that our segmentation performance was superior to other methods.

The remainder of this paper is organized as follows: [Section 2](#) describes our proposed approach in detail. [Section 3](#) presents the fundus dataset, preprocessing methods, and experimental results. Finally, we conclude with a summary and outlook in [Section 4](#).

## 2 Method

### 2.1 Network Architecture

The architecture of the proposed DT-Net is shown in [Fig. 1](#). It consists of four main parts: encoder, decoder, multi-scale input and side output. We improve on U-Net, one of the simplest and most popular architectures in medical image segmentation. Firstly, because the information obtained by the single-scale input of U-Net is limited, we use multi-scale layers to construct image pyramid input, and average pooling is used on the retinal images of size  $H \times W$  to obtain the multi-scale image information, enhancing each layer's feature information. Secondly, a hybrid block is used in the encoder to extract vessel features with irregular shapes and sizes. The encoders are connected through max pooling, which halves the size of the feature map and generates hierarchical features at different levels. Except for the first layer, each layer is inputted by the maximum pool map of the upper layer and the feature map of this layer. The high-level features can correctly identify the coarse vessel information, and the low-level features can accurately obtain the tiny vessel information. Among them, the hybrid block combines modulated deformable convolution and MHSA. The segmentation effect is improved by using deformable convolution for local feature extraction and using MHSA to learn global features. In self-attention, relative position encoding is used to learn the content-position relationship in images.

This paper uses a novel decoder structure to fuse the high-resolution and low-resolution information. The decoder uses dual-input MHSA to obtain low-resolution and high-resolution information and then passes through the residual block [38] to achieve feature reuse, alleviate the problem of gradient disappearance, prevent the occurrence of overfitting, and improve segmentation capabilities. The blue-shaded part at the bottom of [Fig. 1](#) is the structure of the residual block. Finally, multi-scale features are fused using image information at different scales. This structure of first down-sampling and then up-sampling reduces the risk of overfitting to a certain extent. In the side output path, the feature map is spatially up-sampling, and then  $1 \times 1$  convolution is performed to compress the

number of channels to 2, which is convenient for direct comparison with ground truth and outputs the corresponding probability value of each pixel of the image.

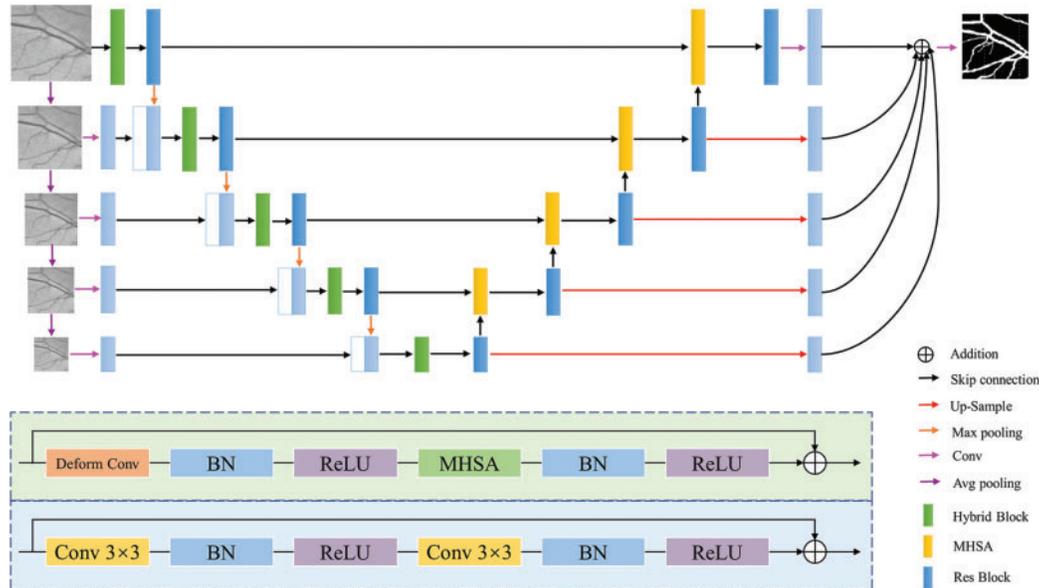


Figure 1: DT-Net network architecture diagram

## 2.2 Deformable Convolution

Most of the original CNN extract feature information at a fixed position in an image based on a fixed receptive field structure, and cannot adaptively generate deformable receptive fields and convolution kernel shapes according to different image features [39]. However, the vessel structure is irregular and complex, and the introduction of deformable convolution can enhance the construction ability of retinal vessel geometric deformation. On the basis of the traditional convolution, the deformable convolution increases the direction vector of the convolution kernel to make the shape of the convolution kernel closer to the feature. A learnable offset is introduced into the deformable convolution. Offset learning is the use of interpolation algorithm, through back propagation learning. The effective receptive field can more accurately cover the actual shape of the vessel to learn more features. Therefore, deformable convolution is used in this paper to enhance the generalization ability of the adaptability to different position information of the image and the mapping ability during the convolution process. The deformable convolution formula is as follows:

$$y(p) = \sum_{i=1}^N w_i \cdot x(p + p_i + \Delta p_i) \cdot \Delta m_i \quad (1)$$

Let  $N$  denotes the sampling position of a given standard convolution kernel, and  $w_i$  and  $p_i$  denote the weight of the  $i$ -th position and the preset offset, respectively.  $x(p)$  and  $y(p)$  denote the features at position  $p$  on the input and output feature maps  $x$  and  $y$ , respectively. Where  $\Delta p_i$  and  $\Delta m_i$  are the learnable offset and adjustment factor at the  $i$ -th position, and the adjustment factor  $\Delta m_i \in [0, 1]$  and  $\Delta p_i$  is an arbitrary value. It can be found that the deformable convolution learns the offset and the weight of the sampling points, which can effectively capture the structural details of tiny vessels and thus achieve more accurate feature extraction.

### 2.3 Multi-Head Self-Attention Mechanism

MHSA is an attention mechanism that pays more attention to the internal structure, inherently has a global receptive field, and is good at capturing long-distance dependencies. The input feature map can be expressed as  $X \in R^{H \times W \times C}$ , where  $H, W, C$  are the height, width, and number of channels, respectively. The self-attention calculation formula is as follows:

$$\text{Attention}(Q, K, V) = \text{softmax}\left(\frac{QK^T}{\sqrt{d}}\right)V = AV \tag{2}$$

where three  $1 \times 1$  convolutions are used to project  $X$  for query, key and value embedding:  $Q, K, V$  in  $R^{H \times W \times d}$ , where  $d$  is the dimension of the embedding for each head. The attention matrix  $A$  works well for feature aggregation, where each row value corresponds to the similarity of a given element in  $Q$  relative to all elements in  $K$ .

Because the image is highly structured data, in the local characteristics of high resolution, in addition to the border area, most of the pixels with similar features. Therefore, computing the attention among all pixels is very inefficient and redundant. So, we propose an efficient self-attention for the task of vessel segmentation, as shown in Fig. 2. The proposed self-attention decoder architecture is used to recover detailed information from the skip connections of the encoder, where  $x$  is the image feature of the previous layer in the decoder, and then the  $1 \times 1$  convolution is performed to obtain a low-resolution image of size  $H_l \times W_l \times d$  characteristics,  $y$  is the image feature from the same layer in the encoder, and then a high-resolution feature of size  $H_h \times W_h \times d$  is obtained by  $1 \times 1$  convolution. Then the dot product and soft-max are performed, and the pairwise attention matrix between the input units. Finally, image features of size  $H_h \times W_h \times d$  are obtained. For positional encoding, standard self-attention blocks lose their positional information and are ineffective for construction highly structured image content [40]. The sinusoidal embedding in the convolution layer in the previous research does not have the property of translation, so the 2-dimensional relative position coding is used by adding the information of relative height  $R_h$  and width  $R_w$ . Relative position coding is used before soft-max operation, and the attention logit is  $qk^T \times qr^T$ .

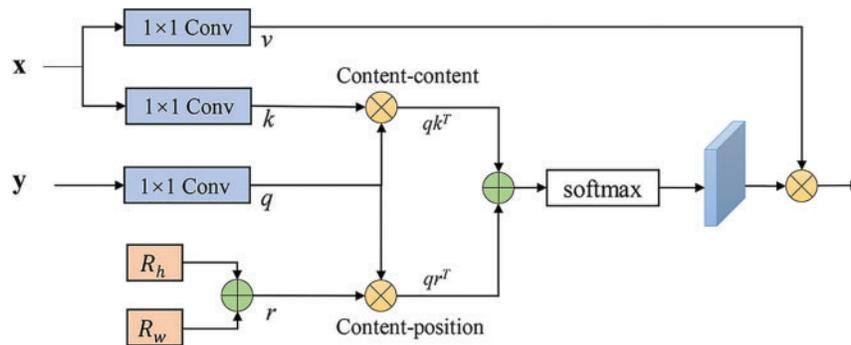


Figure 2: MHSA decoder

### 2.4 Loss Function

Loss function has a significant influence on deep learning training tasks. Most existing methods use only a single loss function to evaluate the network performance. Image segmentation tasks usually use cross-entropy as a loss function, and the ratio of foreground and background pixels in the retinal image is severely imbalanced, resulting in the features of retinal vessels cannot be effectively learned by

the model. In the binary segmentation task, the Dice loss function can alleviate the above problems, and its essence is to measure the degree of overlap between two samples. However, adjusting the weight of the network according to a single loss function can easily lead to the loss of the feature information of the middle and lower layers of the network. Mixed loss can effectively help the model training and enhance segmentation quality. Therefore, the network is trained using a hybrid loss function. Compare the output with the ophthalmologist's criteria and calculate the loss between them:

$$L^{(n)} = \omega \cdot L_{bce} + (1 - \omega) \cdot L_{dice} \quad (3)$$

where  $\omega$  is the weighting factor for balancing different losses. The binary cross entropy (BCE) loss encourages the segmentation model to independently predict the correct class label at each pixel position. Dice loss can alleviate the imbalance of class to some extent. The BCE loss function and Dice loss function are defined as follows:

$$L_{bce} = -\frac{1}{K} \sum_{i=1}^K (g_i \log(p_i) + (1 - g_i) \log(1 - p_i)) \quad (4)$$

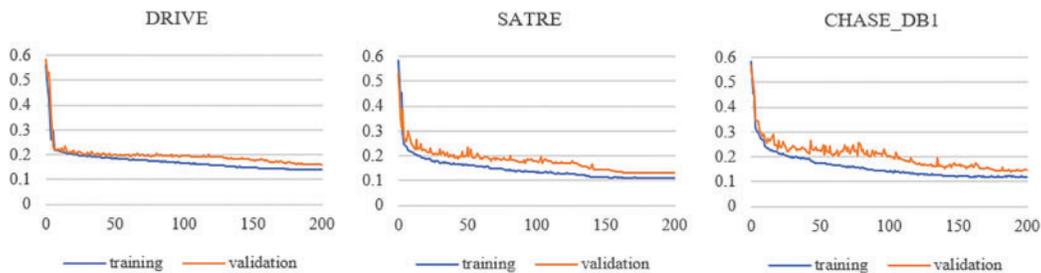
$$L_{dice} = 1 - \frac{2 \sum_{i=1}^K p_i g_i + \epsilon}{\sum_{i=1}^K p_i^2 + \sum_{i=1}^K g_i^2 + \epsilon} \quad (5)$$

where  $K$  represents the number of pixels in a given image, and  $p_i \in [0, 1]$ ,  $g_i \in [0, 1]$  represent the predicted probability and label probability of the  $i$ -th pixel, respectively. The parameter  $\epsilon$  is a Laplace smoothing factor, which avoids numerical problems and speeds up the convergence of the training process.

### 3 Experimental

#### 3.1 Experimental Details

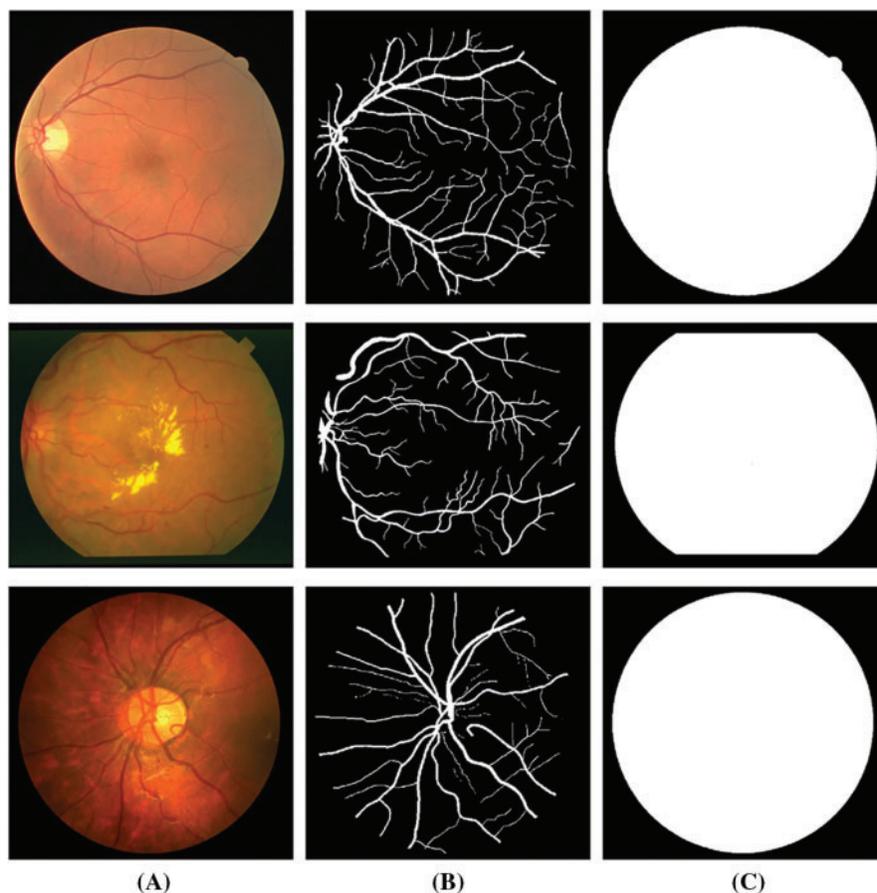
In this paper, we run all experiments based on Windows 10, Intel Core i5-10400F CPU, GeForce GTX 1080ti GPU, Python 3.7 language, and PyTorch deep learning framework. The parameters in the network are optimized using the Adam optimizer with an initial learning rate of 0.0005 and a weight decay of 0.001. To dynamically adjust the training process, a cosine annealing strategy is utilized to update the learning rate. The proposed DT-Net framework is trained for 200 epochs with a batch size of 2. Fig. 3 shows the loss function curves of the proposed method for training and verifying datasets relative to iteration on three datasets: DRIVE [41], STARE [42], and CHASE\_DB1 [43]. The horizontal coordinate of the image is the iteration period "Epoch", and the ordinate is the loss value "Loss". Legend "training" means training, and Legend "validation" means validation. When the proposed method is trained on three datasets, the training and validation losses converge rapidly within 50 epochs, flatten out within 150 epochs, and then reach a stable value.



**Figure 3:** Loss function curves vs. iterations for the training and validation datasets

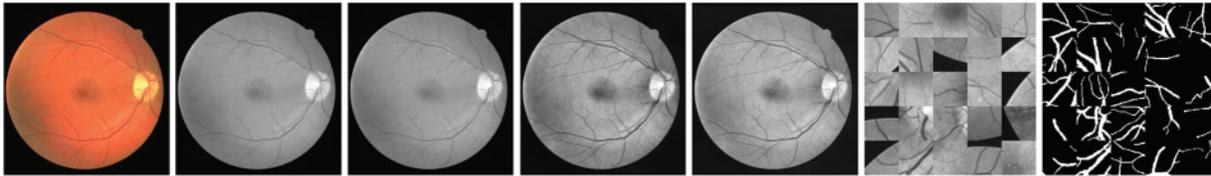
### 3.2 Data and Preprocessing

We use the public and widely used standard datasets DRIVE, STARE, and CHASE\_DB1 as the training and testing datasets of the proposed network. Sample images for the three datasets are shown in Fig. 4, along with their ground truth vessel segmentation masks and field-of-view (FOV) masks. The DRIVE dataset consists of 40 fundus images with an image resolution of  $584 \times 565$ , and it is specified that the training set and the test set both contain 20 images. The STARE dataset consists of 20 fundus images with an image resolution of  $700 \times 605$ , including 10 retinal images with pathological features. This dataset can evaluate the ability of the model to segment abnormal fundus images. The CHASE\_DB1 dataset consists of 28 retinal images of 14 children with an image resolution of  $960 \times 999$ . Since the STARE and CHASE\_DB1 datasets do not officially specify the training set and test set, we use the first 10 images of the STARE dataset for model training according to DUNet [29], and the remaining 10 images are used for model performance evaluation. For the CHASE\_DB1 datasets, we follow a common protocol [44], selecting the first 20 images for model training and the remaining 8 images for model performance evaluation. The three datasets contain the results of the manual segmentation of retinal images by two experienced ophthalmologists. We used the segmentation results of the first ophthalmologist as the ground truth for network training [45] and also as the standard segmentation results for network model and algorithm evaluation.



**Figure 4:** Sample images from DRIVE, STARE, and CHASE\_DB1. (A) Original image; (B) ground truth; (C) field-of-view masks

Retinal images often contain noise and uneven illumination, so all images from these three datasets undergo four kinds of preprocessing for image enhancement before the training and testing datasets of the network. The preprocessing process of the fundus images is shown in Fig. 5. Firstly, the color images are converted into grayscale images, simplifying the subsequent preprocessing steps and reducing the computation during training. Secondly, each pixel in the grayscale image is normalized to reduce the data dimension and speed up the convergence. Then, the CLAHE method [46] is used to suppress image noise and enhance the details of the vessel and the contrast between the vessel and background. Finally, nonlinear transformation and gamma correction are performed to solve the image quality problem caused by the brightness of the input images, enhance the contrast, make the vessel in the darker area clearer, and improve the image quality. After the above processing, it can be found that the distinction between the retinal vessel and the background is significantly improved at this time, which is conducive to feature extraction in the training process and enhances the segmentation quality of the retinal vessel.



**Figure 5:** Image preprocessing of fundus images, from left to right, are original image, image graying, image normalization, gradient histogram equalization, gamma correction, original images patches, and ground truth patches

Due to the limited number of images in the fundus datasets, patch processing is adopted to expand the datasets to reduce the effect of overfitting. In the training process, each image of the pre-processed datasets is first randomly cropped into a patch of size  $64 \times 64$  for the training of the network. The corresponding patches are also extracted from the ground truth to ensure the original images and the ground truth. In the experiments, 90% of the extracted patches are used for training and the remaining 10% are used for validation. Fig. 5 shows some image patches and the corresponding ground truth of the fundus images.

### 3.3 Evaluation Index

Similar to most methods of retinal image segmentation, we will compare the proposed DT-Net method with other algorithms and evaluate it through the following indicators: Accuracy (ACC), Specificity (SP), Sensitivity (SE), F1-score, and Area Under Receiver Operating Characteristic (ROC) Curve (AUC). Acc is used to evaluate the overall segmentation performance of the model. The larger the ACC, the more accurate the segmentation. The specific mathematical expression is as follows:

$$ACC = \frac{TP + TN}{TP + FN + FP + TN} \quad (6)$$

SP is an important metric of retinal vessel segmentation. It is the ratio of correct negative predictions to the total number of negative predictions. It mainly evaluates the ability to recognize background in retinal images. The better the SP value, the lower the false positive rate (FPR). The specific mathematical expression is as follows:

$$SP = \frac{TN}{TN + FP} \quad (7)$$

SE mainly evaluates the ability to recognize retinal vessels (positive) in retinal images. It is the ratio of correct positive predictions to the total number of positive predictions in the predicted results. The specific mathematical expression is as follows:

$$SE = \frac{TP}{TP + FN} \quad (8)$$

F1-score for evaluation of segmentation results and the similarity criteria of ophthalmology. The larger the value is, the closer the algorithm segmentation result is to the gold standard, and the segmentation effect. The specific mathematical expression is as follows:

$$F1 = \frac{2TP}{2TP + FP + FN} \quad (9)$$

Among them, true positive (TP) means correctly identified vessel pixels, true negative (TN) means correctly identified non-vessel pixels, false positive (FP) means non-vessel pixels identified as vessels, and false negative (FN) means vessel pixels identified as non-vessel.

In addition, we also introduce AUC to evaluate the segmentation performance of the model. AUC is a professional metric of retinal vessel segmentation. The closer its value is to 1, the better the performance of the algorithm and the stronger the robustness. The ROC curve describes the relationship between the true position rate and the false position rate under the different classification thresholds. The closer the value of the area under the AUC is to 1, the better the algorithm performs and the more robustness.

### 3.4 Ablation Experiment

To further verify the effectiveness of the proposed network for vessel segmentation, we conduct ablation experiments on the DRIVE dataset. The prediction results of the network are compared in terms of five performance metrics: ACC, SE, SP, AUC, and F1-score. To more clearly see the improvement of the accuracy of retinal vessel segmentation by each module proposed in the model, the segmentation performance of different methods is shown in Table 1. M0 uses a hybrid loss function based on U-Net. M1 adds multiple inputs and side outputs based on M0. M2 adds the encoder hybrid block based on M1. M3 adds the transformer decoder block based on M1. M4 adds the encoder hybrid block and transformer decoder block based on M1.

**Table 1:** Ablation experimental results of vessel segmentation on DRIVE dataset

Method	ACC	SE	SP	AUC	F1-score
M0	95.58%	74.68%	98.62%	96.70%	81.13%
M1	95.63%	81.24%	97.73%	97.89%	82.56%
M2	96.25%	83.75%	97.99%	98.40%	84.57%
M3	96.26%	85.49%	97.76%	98.19%	84.86%
M4	<b>96.31%</b>	<b>86.36%</b>	<b>98.84%</b>	<b>98.43%</b>	<b>84.88%</b>

As shown in Table 1, when multi-scale input and side output is added to M0, each index is significantly improved, and the segmentation performance of the network is improved. After adding the hybrid block to the network, AUC and F1-score in M2 are 0.51% and 2.01% higher than M1, respectively, which verifies the effectiveness of the hybrid block. In M3, SE, AUC, and F1-score

are 4.25%, 0.3%, and 2.3% higher than M1, respectively. It is shown that the proposed MHSA decoder block is effective in retinal vessel segmentation and enhances the performance of retinal vessel segmentation.

We can see from the last row of [Table 1](#) that the values of SE, AUC, and F1-score of the proposed network are increased from 74.68%, 96.70%, 81.13% of M0 to 86.36%, 98.43%, 84.88%, respectively. Experiments show that using either the hybrid block of the encoder or the attention block of the decoder can improve the segmentation performance of the network, which shows their rationality and effectiveness. Therefore, the proposed method has advantages in retinal vessel segmentation.

Ablation experiments are performed by setting different loss functions to verify which loss function is more suitable for the proposed method. The effects of varying loss functions on performance indexes are shown in [Table 2](#). First, “DT+BCE” uses a BCE loss function to train the network. “DT+Dice” uses the Dice loss function. “DT+BCE+Dice” combines the BCE loss function and the Dice loss function. The results in [Table 2](#) show that almost all the metrics are improved with the help of the hybrid loss, which proves that the hybrid loss contributes to enhancing the accuracy of the model.

**Table 2:** Loss function ablation experiment

Method	ACC	SE	SP	AUC	F1-score
DT+BCE	96.22%	85.88%	97.67%	98.18%	84.79%
DT+Dice	96.21%	86.28%	97.58%	96.62%	84.82%
DT+BCE+Dice	<b>96.31%</b>	<b>86.36%</b>	<b>98.84%</b>	<b>98.43%</b>	<b>84.88%</b>

The learning rate, as an essential parameter in the process of model training, controls the learning progress of the network model. To explore the influence of different learning rates on segmented images, [Table 3](#) shows the segmentation results when the learning rate of this method is 0.0001, 0.0003, 0.0005, 0.0007, 0.0009, and 0.0011. When the learning rate is set to 0.0005, the best performance is achieved for all metrics. When the learning rate increases or decreases, both F1-score and AUC will decrease.

**Table 3:** Segmentation results with different learning rates

learning rate	ACC	SE	SP	AUC	F1-score
0.0001	96.18%	85.14%	97.73%	98.19%	84.56%
0.0003	96.20%	<b>86.39%</b>	97.65%	98.11%	84.73%
<b>0.0005</b>	<b>96.31%</b>	86.36%	<b>98.84%</b>	<b>98.43%</b>	<b>84.88%</b>
0.0007	96.15%	85.58%	97.62%	98.09%	84.01%
0.0009	95.66%	85.88%	97.19%	98.21%	83.59%
0.0011	96.28%	87.74%	98.03%	98.08%	83.81%

### 3.5 Comparison with Other Methods

We chose five other retinal vessel segmentation methods to compare with our method to prove the advantages of the proposed method. The five methods are U-Net [18], DUNet [24], GT U-Net [47], Attention Residual U Network (AReN-U-Net) [48], and Multistage Dual-Path Interactive

Refinement Network (Multistage DPIRef-Net) [49]. For a better comparison with these methods, these models are trained using the same experimental settings as in this paper. In addition, we compare the performance with UNet++ [27], Li et al. proposed method [44], D-MNet [50], WA-Net [51], and TUNet-LBF [52]. Among them, GT U-Net modifies both the encoder and decoder to achieve good performance on tooth segmentation. The AReN-UNet proposes a novel cascaded network driven by integrating attention and residual modules. It improves convergence and stability and reduces vessel breakdown in the vessel map. Multistage DPIRef-Net uses multi-stage dual-path interaction to refine the network, retain the vascular branch edges, suppress the false positive rate, and accurately segment the arteriovenous vascular map of the retinal surface. The Multistage DPIRef-Net is a vessel segmentation architecture with a single encoder and dual decoder. It requires annotated labels of arteries and veins, while our network uses only one kind of annotated label. Therefore, to be fair, we use the experimental data of this paper in the same experimental setting, i.e., using one kind of annotated label. The modified network mainly consists of one backbone network and three single-path stages. The D-MNet uses deformable convolution and attention modules to improve the accuracy and connectivity of vessel segmentation.

Tables 4–6 evaluate the different vessel segmentation methods in the DRIVE, STARE, and CHASE\_DB1 datasets. Because there are more background pixels than vessel pixels in the fundus images, AUC and F1-score metrics are more suitable for evaluating the vessel segmentation method. In Table 4, compared with the maximum of existing methods, our proposed method performs better on the DRIVE dataset. There is a 2.83% increase in SE, 0.2% in SP, 0.22% in AUC, and 1.22% in F1-score, respectively. The highest SE and SP of the proposed model means that retinal vessels can be identified more accurately, and noise information can be suppressed. This is because the MHSA mechanism focuses on capturing global vessel details. We can observe from Table 5 that the proposed method achieves the best performance for ACC, SP, AUC, and F1-score on STARE datasets compared to other methods. The proposed method performs better, indicating that the framework is effective for vessel segmentation. Since there are many lesion images in the STARE dataset, the SE is not optimal all metrics of our method are highest on the CHASE\_DB1 dataset except the SE and ACC metric. On the whole, our method performs well. As seen from Tables 4–6, the proposed method has the highest F1-score metrics on the three datasets compared to the maximum value of each metric of the other methods, with an increase of 1.85%, 3.61%, and 0.26%, respectively. This indicates that the proposed method can distinguish retinal vessel pixels and background pixels effectively and accurately. In general, compared with these methods, the proposed method can segment retinal vessels more accurately and has good prospects for application in clinical medical imaging diagnosis.

**Table 4:** Comparison of the proposed method with existing methods in the DRIVE dataset

Method	Year	ACC	SE	SP	AUC	F1-score
U-Net	2015	95.58%	74.91%	98.59%	96.83%	81.19%
DU-Net	2019	95.49%	75.84%	98.35%	97.59%	81.05%
UNet++	2020	95.35%	74.73%	98.35%	97.13%	80.35%
GT U-Net	2021	96.22%	80.58%	98.64%	97.96%	83.66%
AReN-UNet	2021	<b>96.60%</b>	83.53%	98.12%	98.21%	82.87%
Li et al.	2021	95.68%	79.21%	98.10%	98.06%	–
Multistage DPIRef-Net	2022	95.64%	80.53%	98.39%	97.62%	81.61%

(Continued)

**Table 4 (continued)**

Method	Year	ACC	SE	SP	AUC	F1-score
D-MNet	2022	95.39%	73.68%	97.12%	97.93%	82.13%
WA-Net	2022	95.75%	79.66%	98.10%	97.84%	82.69%
TUnet_LBF	2023	96.50%	81.40%	98.09%	89.75%	–
Proposed	2023	96.31%	<b>86.36%</b>	<b>98.84%</b>	<b>98.43%</b>	<b>84.88%</b>

**Table 5:** Comparison of the proposed method with existing methods in the STARE dataset

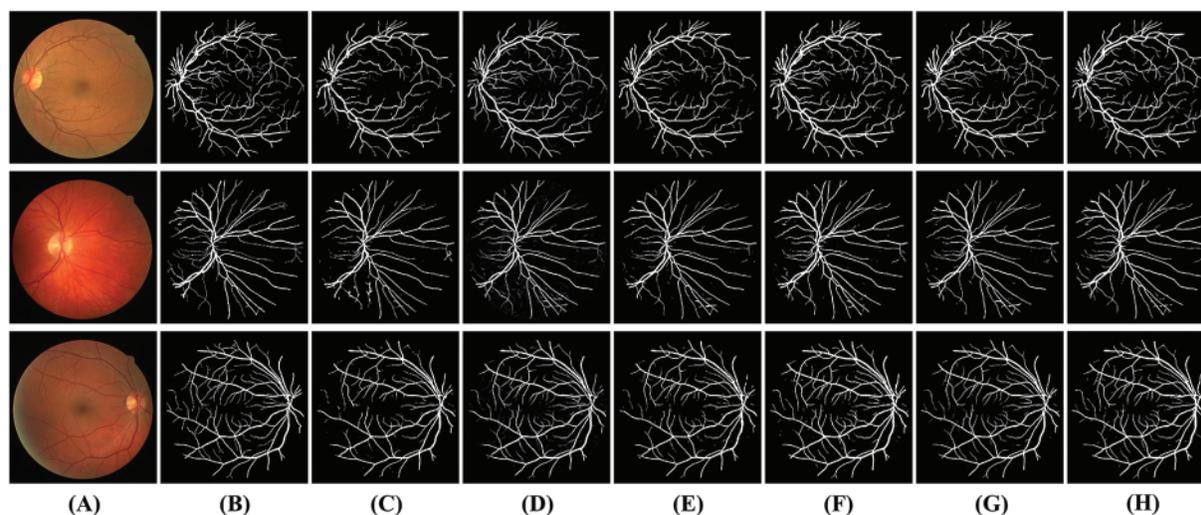
Method	Year	ACC	SE	SP	AUC	F1-score
U-Net	2015	95.81%	77.62%	97.90%	97.12%	79.28%
DU-Net	2019	96.15%	71.43%	99.04%	97.23%	76.55%
UNet++	2020	96.05%	77.76%	98.32%	97.40%	81.32%
GT U-Net	2021	95.87%	70.49%	98.49%	97.03%	77.81%
AReN-UNet	2021	96.73%	81.15%	98.78%	98.59%	82.11%
Li et al.	2021	96.78%	83.52%	98.23%	98.75%	–
Multistage DPIRef-Net	2022	96.59%	80.04%	98.55%	97.64%	80.66%
D-MNet	2022	96.43%	<b>84.35%</b>	97.79%	98.55%	82.74%
WA-Net	2022	96.55%	77.67%	98.77%	96.65%	81.76%
TUnet_LBF	2023	96.81%	80.04%	98.52%	89.28%	–
Proposed	2023	<b>97.03%</b>	83.52%	<b>99.37%</b>	<b>99.00%</b>	<b>86.37%</b>

**Table 6:** Comparison of the proposed method with existing methods in the CHASE\_DB1 dataset

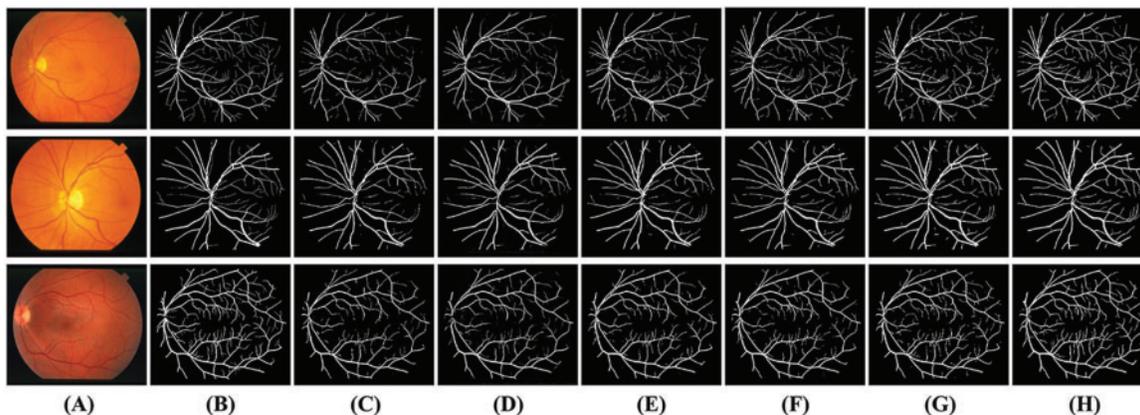
Method	Year	ACC	SE	SP	AUC	F1-score
U-Net	2015	95.96%	73.38%	97.72%	96.54%	79.10%
DU-Net	2019	96.69%	81.53%	98.30%	98.40%	80.43%
UNet++	2020	95.06%	63.61%	98.94%	97.04%	73.90%
GT U-Net	2021	95.96%	72.72%	99.05%	96.11%	80.27%
AReN-UNet	2021	97.51%	83.18%	98.41%	98.45%	81.95%
Li et al.	2021	96.35%	78.18%	98.19%	98.10%	–
Multistage DPIRef-Net	2022	96.45%	81.18%	98.52%	97.79%	80.75%
D-MNet	2022	95.87%	<b>85.43%</b>	96.93%	98.06%	79.01%
WA-Net	2022	<b>98.41%</b>	80.42%	98.26%	96.53%	80.98%
TUnet_LBF	2023	97.08%	83.54%	98.06%	90.80%	–
Proposed	2023	97.37%	84.87%	<b>99.21%</b>	<b>98.48%</b>	<b>82.74%</b>

To further observe the segmentation results of the models, partial segmentation results on the three datasets are given for visual comparison, as shown in Figs. 6–8. It can be seen that the DT-Net model

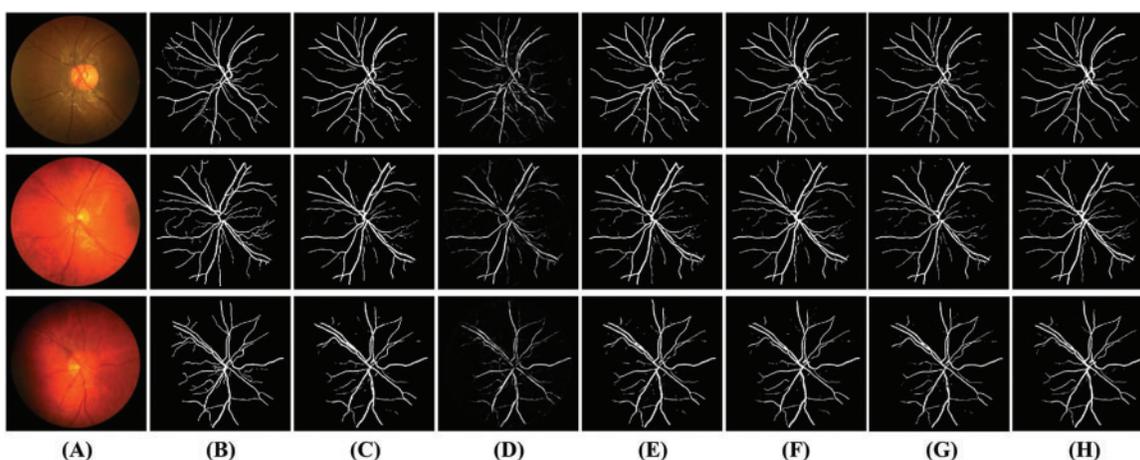
produces more details of the vessel segmentation. Compared with U-Net, DT-Net can detect more vessels. Compared with DUNet and GT U-Net, DT-Net can detect some details of missing vessels and thus complete segmentation more efficiently. Compared with AReN-UNet and Multistage DPIRef-Net, DT-Net has better vessel continuity. We can observe that the DT-Net is superior to the other five methods on three datasets. As can be seen in Figs. 6–8, the segmentation effect of the proposed method is very close to the standard of ophthalmologist manual segmentation, obtaining more continuous vessels. It can successfully segment the continuous tiny vessels and has good generalization ability when segmenting different datasets. This proves the network can reduce background noise, enhance contrast, and preserve irregular vessels well. Visualization further illustrates the importance of multi-scale contextual information and capturing long-term dependencies in retinal vessel segmentation. This suggests that the proposed method can segment retinal vessel images, help specialized physicians in disease diagnosis, and reduce the workload of clinical medical specialists. In addition, we use the ROC curve to evaluate the model, shown in Fig. 9. The closer the ROC curve is to the upper left boundary, the more accurate the network is. Fig. 10 shows a locally enlarged view of the tiny vessel in the segmentation result. This is because, in retinal images, tiny vessels are not significantly different from the image background. Therefore, to help the network pay attention to essential features and suppress unnecessary features, we use the MHSA mechanism in both the encoder and decoder. It can be seen from Fig. 10 that the proposed algorithm has good robustness to the intersection of the vessel and tiny vessel areas with low contrast and maintains the degree and connectivity of thick and thin vessels, and the segmentation results of the lesion region are relatively close to the standard segmentation. The reliability and robustness of this algorithm for retinal vessel segmentation are verified. The above experimental results can prove that the performance of the proposed model is generally better, it can more accurately identify vessels and backgrounds, and segment tiny vessels better.



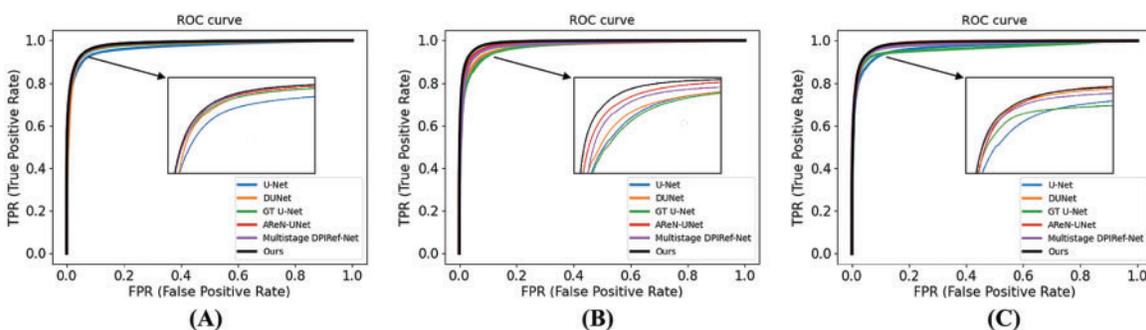
**Figure 6:** The segmentation results of different models on DRIVE datasets. (A) Original images; (B) ground truth images; (C) U-Net; (D) DUNet; (E) GT U-Net; (F) AReN-UNet; (G) multistage DPIRef-Net; (H) ours



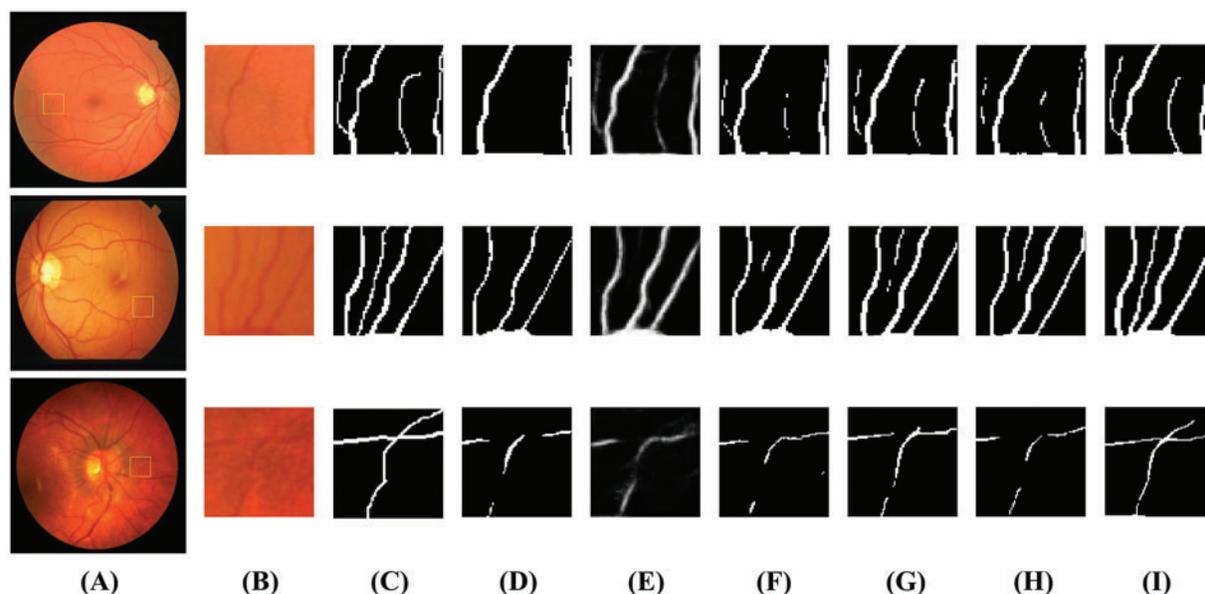
**Figure 7:** The segmentation results of different models on STARE datasets. (A) Original images; (B) ground truth images; (C) U-Net; (D) DUNet; (E) GT U-Net; (F) AReN-UNet; (G) multistage DPIRef-Net; (H) ours



**Figure 8:** The segmentation results of different models on CHASE\_DB1 datasets. (A) Original images; (B) ground truth images; (C) U-Net; (D) DUNet; (E) GT U-Net; (F) AReN-UNet; (G) multistage DPIRef-Net; (H) ours



**Figure 9:** The ROC curves of the DT-Net model on different datasets. (A) DRIVE dataset. (B) STARE dataset. (C) CHASE DB1 dataset



**Figure 10:** Partial increased view of different models on different datasets. From top to bottom are the fundus images from the DRIVE, STARE, and CHASE DB1 datasets. From left to right: (A) original images; (B) partial views; (C) ground truth; (D) U-Net; (E) DUNet; (F) GT U-Net; (G) AREN-UNet; (H) multistage DPIRef-Net; (I) ours

#### 4 Discussion

In our work, we propose a hybrid convolution and transformer network evolved from the classical model U-Net, which aims to aggregate multi-scale feature information at different resolutions to achieve accurate and efficient vessel segmentation. The fundus image is full of noise and low contrast. Therefore, we first preprocess to improve image contrast and suppress the background noise of the source image. To fully use multi-scale information, DT-Net uses multi-scale images as input, and then we introduce deformable convolution to change the convolution kernel according to the actual shape of blood vessels to obtain more accurate structural information. Meanwhile, the MHSA mechanism is used to capture the distant relationship of fundus images, making up for the defect that CNN cannot extract global features.

In addition, the proposed network is verified by ablative experiments. The ACC and AUC indexes of the network improved significantly after the addition of a mixing block to the encoder, and the SE and F1-scoring indexes of the network improved significantly after the addition of a transformer decoder block. Of course, the current study of DT-Net proposed has the following shortcomings: (1) Due to the similarity between the background and blood vessels in the datasets, our method cannot achieve the best performance in each index; (2) They inevitably lose some container details due to constant up-sampling. In the future, we will introduce more advanced methods, such as the encoding pattern in Swin-UNet, to preserve more details in the original image and make our model perform better on various metrics.

## 5 Conclusion

We propose a network named DT-Net for fundus blood vessel segmentation. The performance of this method is mainly improved by the introduction of variable convolution and multiple self-attention mechanisms, which not only extract the structural information easily ignored in fundus blood vessel images but also effectively extract information at different scales. And the DT-Net presented in the DRIVE, STARE, and CHASE\_DB1 datasets is significantly improved. Experimental results show that this method can better process different fundus data sets, has better generalization ability, and provides more accurate segmentation results for medical diagnosis and treatment. In terms of segmentation results, our model can segment more vascular details and has better connectivity.

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**Author Contributions:** WJ and SM design of study, analysis of data. WJ and SM conducted experiments and drafted the manuscript. PG and SM revised and edited the manuscript. YS and PG polished the manuscript. All authors reviewed the results and approved the final version of the manuscript.

**Availability of Data and Materials:** Publicly available datasets were analyzed in this study. This data can be found here: DRIVE: <https://drive.grand-challenge.org>; STARE: <https://cecas.clemson.edu/~ahoover/stare/>; CHASE\_DB1: <https://blogs.kingston.ac.uk/retinal/chasedb1>.

**Conflicts of Interest:** The authors declare that they have no conflicts of interest to report regarding the present study.

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